RAW SEQUENCE LISTING   DATE: 07/16/2003
PATENT APPLICATION: **US/09/866,248C**   TIME: 08:05:33

Input Set : **A:\57155A.txt**
Output Set: **N:\CRF4\07162003\I866248C.raw**

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   5 <110> APPLICANT: Gerald, Christophe P.G.
   7         Jones, Kenneth A.
   9         Bonini, James A.
  11         Borowsky, Beth
  15 <120> TITLE OF INVENTION: DNA Encoding Mammalian Neuropeptide FF (NPFF) Receptors
  17       and Uses Thereof
  21 <130> FILE REFERENCE: 1795/57155-A
C--> 25 <140> CURRENT APPLICATION NUMBER: US/09/866,248C
C--> 27 <141> CURRENT FILING DATE: 2001-05-25
  31 <150> PRIOR APPLICATION NUMBER: 09/161,113
  33 <151> PRIOR FILING DATE: 1998-09-25
  37 <160> NUMBER OF SEQ ID NOS: 42
  41 <170> SOFTWARE: PatentIn Ver. 2.0 - beta
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  47 <211> LENGTH: 1410
  49 <212> TYPE: DNA
  51 <213> ORGANISM: Rattus norvegicus
  55 <400> SEQUENCE: 1
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  59 ggagcagaca gtatggaggc ggagccctcc cagcctccca acggcagctg gcccctgggt 120
  61 cagaacggga gtgatgtgga gaccagcatg gcaaccagcc tcaccttctc ctcctactac 180
  63 caacactcct ctccggtggc agccatgttc atcgcggcct acgtgctcat cttcctcctc 240
  65 tgcatggtgg gcaacaccct ggtctgcttc attgtgctca agaaccggca catgcgcact 300
  67 gtcaccaaca tgtttatcct caacctggcc gtcagcgacc tgctggtggg catcttctgc 360
  69 atgcccacaa cccttgtgga caaccttatc actggttggc cttttgacaa cgccacatgc 420
  71 aagatgagcg gcttggtgca gggcatgtcc gtgtctgcat cggtttttcac actggtggcc 480
  73 atcgctgtgg aaaggttccg ctgcatcgtg cacccttttcc gcgagaagct gacccttcgg 540
  75 aaggcgctgt tcaccatcgc ggtgatctgg gctctggcgc tgctcatcat gtgtccctcg 600
  77 gcggtcactc tgacagtcac ccgagaggag catcacttca tgctggatgc tcgtaaccgc 660
  79 tcctacccgc tctactcgtg ctgggaggcc tggcccgaga agggcatgcg caaggtctac 720
  81 accgcggtgc tcttcgcgca tctctacctg gtgccgctgg cgctcatcgt agtgatgtac 780
  83 gtgcgcatcg cgcgcaagct atgccaggcc cccggtcctg cgcgcgacac ggaggaggcg 840
  85 gtggccgagg gtggccgcac ttcgcgccgt agggcccgcg tggtgcacat gctggtcatg 900
  87 gtggcgctct tcttcacgtt gtcctggctg ccactctggg tgctgctgct gctcatcgac 960
  89 tatggggagc tgagcgagct gcaactgcac ctgctgtcgg tctacgcctt ccccttggca 1020
  91 cactggctgg ccttcttcca cagcagcgcc aacccccatca tctacggcta cttcaacgag 1080
  93 aacttccgcc gcggcttcca ggctgccttc cgtgcacagc tctgctggcc tccctgggcc 1140
  95 gcccacaagc aagcctactc ggagcggccc aaccgcctcc tgcgcaggcg ggtggtggtg 1200
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  99 gggcctggcc ggctgccact gcgcaatggg cgtgtggccc atcaggatgg cccggggggaa 1320
 101 gggccaggct gcaaccacat gcccctcacc atcccggcct ggaacatttg aggtggtcca 1380
 103 gagaagggag ggccagtagt cctgtggccc                                  1410
 107 <210> SEQ ID NO: 2
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RAW SEQUENCE LISTING   DATE: 07/16/2003
PATENT APPLICATION: **US/09/866,248C**   TIME: 08:05:33

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Output Set: **N:\CRF4\07162003\I866248C.raw**

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113 <213> ORGANISM: Rattus norvegicus
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127              20                  25                  30
131 Ser Ser Tyr Tyr Gln His Ser Ser Pro Val Ala Ala Met Phe Ile Ala
133            35                  40                  45
137 Ala Tyr Val Leu Ile Phe Leu Leu Cys Met Val Gly Asn Thr Leu Val
139       50                  55                  60
143 Cys Phe Ile Val Leu Lys Asn Arg His Met Arg Thr Val Thr Asn Met
145   65                  70                  75                  80
149 Phe Ile Leu Asn Leu Ala Val Ser Asp Leu Leu Val Gly Ile Phe Cys
151                  85                  90                  95
155 Met Pro Thr Thr Leu Val Asp Asn Leu Ile Thr Gly Trp Pro Phe Asp
157              100                 105                 110
161 Asn Ala Thr Cys Lys Met Ser Gly Leu Val Gln Gly Met Ser Val Ser
163            115                 120                 125
167 Ala Ser Val Phe Thr Leu Val Ala Ile Ala Val Glu Arg Phe Arg Cys
169       130                 135                 140
173 Ile Val His Pro Phe Arg Glu Lys Leu Thr Leu Arg Lys Ala Leu Phe
175 145                 150                 155                 160
179 Thr Ile Ala Val Ile Trp Ala Leu Ala Leu Leu Ile Met Cys Pro Ser
181              165                 170                 175
185 Ala Val Thr Leu Thr Val Thr Arg Glu Glu His His Phe Met Leu Asp
187              180                 185                 190
191 Ala Arg Asn Arg Ser Tyr Pro Leu Tyr Ser Cys Trp Glu Ala Trp Pro
193            195                 200                 205
197 Glu Lys Gly Met Arg Lys Val Tyr Thr Ala Val Leu Phe Ala His Ile
199       210                 215                 220
203 Tyr Leu Val Pro Leu Ala Leu Ile Val Val Met Tyr Val Arg Ile Ala
205 225                 230                 235                 240
209 Arg Lys Leu Cys Gln Ala Pro Gly Pro Ala Arg Asp Thr Glu Glu Ala
211              245                 250                 255
215 Val Ala Glu Gly Gly Arg Thr Ser Arg Arg Arg Ala Arg Val Val His
217              260                 265                 270
221 Met Leu Val Met Val Ala Leu Phe Phe Thr Leu Ser Trp Leu Pro Leu
223            275                 280                 285
227 Trp Val Leu Leu Leu Leu Ile Asp Tyr Gly Glu Leu Ser Glu Leu Gln
229       290                 295                 300
233 Leu His Leu Leu Ser Val Tyr Ala Phe Pro Leu Ala His Trp Leu Ala
235 305                 310                 315                 320
239 Phe Phe His Ser Ser Ala Asn Pro Ile Ile Tyr Gly Tyr Phe Asn Glu
241              325                 330                 335
245 Asn Phe Arg Arg Gly Phe Gln Ala Ala Phe Arg Ala Gln Leu Cys Trp
247            340                 345                 350
251 Pro Pro Trp Ala Ala His Lys Gln Ala Tyr Ser Glu Arg Pro Asn Arg
```

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253         355                   360                   365
257 Leu Leu Arg Arg Arg Val Val Val Asp Val Gln Pro Ser Asp Ser Gly
259     370           375               380
263 Leu Pro Ser Glu Ser Gly Pro Ser Ser Gly Val Pro Gly Pro Gly Arg
265 385               390               395               400
269 Leu Pro Leu Arg Asn Gly Arg Val Ala His Gln Asp Gly Pro Gly Glu
271             405               410               415
275 Gly Pro Gly Cys Asn His Met Pro Leu Thr Ile Pro Ala Trp Asn Ile
277             420               425               430
289 <210> SEQ ID NO: 3
291 <211> LENGTH: 200
293 <212> TYPE: DNA
295 <213> ORGANISM: Homo sapiens
299 <400> SEQUENCE: 3
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303 gccaccccgg ctacaaacct caccttctcc tcctactatc agcacacctc ccctgtggcg 120
305 gccatgttca ttgtggccta tgcgctcatc ttcctgctct gcatggtggg caacaccctg 180
307 gtctgtttca tcgtgctcaa                                           200
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313 <211> LENGTH: 66
315 <212> TYPE: PRT
317 <213> ORGANISM: Homo sapiens
321 <400> SEQUENCE: 4
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325   1               5                   10                  15
329 Thr Asn Thr Glu Ala Thr Pro Ala Thr Asn Leu Thr Phe Ser Ser Tyr
331             20                25                30
335 Tyr Gln His Thr Ser Pro Val Ala Ala Met Phe Ile Val Ala Tyr Ala
337         35                40                45
341 Leu Ile Phe Leu Leu Cys Met Val Gly Asn Thr Leu Val Cys Phe Ile
343     50                55                60
347 Val Leu
349   65
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357 <211> LENGTH: 1302
359 <212> TYPE: DNA
361 <213> ORGANISM: Homo sapiens
365 <400> SEQUENCE: 5
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369 gaaaactggc atcccatctg gaatgtcaat gacacaaagc atcatctgta ctcagatatt 120
371 aatattacct atgtgaacta ctatcttcac cagcctcaag tggcagcaat cttcattatt 180
373 tcctactttc tgatcttctt tttgtgcatg atgggaaata ctgtggtttg ctttattgta 240
375 atgaggaaca acatatgca cacagtcact aatctcttca tcttaaacct ggccataagt 300
377 gatttactag ttggcatatt ctgcatgcct ataacactgc tggacaatat tatagcagga 360
379 tggccatttg gaaacacgat gtgcaagatc agtggattgg tccagggaat atctgtcgca 420
381 gcttcagtct ttacgttagt tgcaattgct gtagataggt tccagtgtgt ggtctaccct 480
383 tttaaaccaa agctcactat caagacagcg tttgtcatta ttatgatcat ctgggtccta 540
385 gccatcacca ttatgtctcc atctgcagta atgttacatg tgcaagaaga aaaatattac 600
387 cgagtgagac tcaactccca gaataaaacc agtccagtct actggtgccg ggaagactgg 660
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Input Set : **A:\57155A.txt**
Output Set: **N:\CRF4\07162003\I866248C.raw**

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391 cccctctccc tcattgtcat catgtatgga aggattggaa tttcactctt cagggctgca 780
393 gttcctcaca caggcaggaa gaaccaggag cagtggcacg tggtgtccag gaagaagcag 840
395 aagatcatta agatgctcct gattgtggcc ctgctttta ttctctcatg gctgcccctg 900
397 tggactctaa tgatgctctc agactacgct gacctttctc caaatgaact gcagatcatc 960
399 aacatctaca tctacccttt tgcacactgg ctggcattcg gcaacagcag tgtcaatccc 1020
401 atcatttatg gtttcttcaa cgagaatttc cgccgtggtt tccaagaagc tttccagctc 1080
403 cagctctgcc aaaaaagagc aaagcctatg gaagcttatg ccctaaaagc taaaagccat 1140
405 gtgctcataa acacatctaa tcagcttgtc caggaatcta catttcaaaa ccctcatggg 1200
407 gaaaccttgc tttataggaa aagtgctgaa aaaccccaac aggaattagt gatggaagaa 1260
409 ttaaaagaaa ctactaacag cagtgagatt taaaaagagc ta                      1302
413 <210> SEQ ID NO: 6
415 <211> LENGTH: 420
417 <212> TYPE: PRT
419 <213> ORGANISM: Homo sapiens
423 <400> SEQUENCE: 6
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431 Trp Asn Val Asn Asp Thr Lys His His Leu Tyr Ser Asp Ile Asn Ile
433              20                  25                  30
437 Thr Tyr Val Asn Tyr Tyr Leu His Gln Pro Gln Val Ala Ala Ile Phe
439              35                  40                  45
443 Ile Ile Ser Tyr Phe Leu Ile Phe Phe Leu Cys Met Met Gly Asn Thr
445          50                  55                  60
449 Val Val Cys Phe Ile Val Met Arg Asn Lys His Met His Thr Val Thr
451  65                  70                  75                  80
455 Asn Leu Phe Ile Leu Asn Leu Ala Ile Ser Asp Leu Leu Val Gly Ile
457              85                  90                  95
461 Phe Cys Met Pro Ile Thr Leu Leu Asp Asn Ile Ile Ala Gly Trp Pro
463              100                 105                 110
467 Phe Gly Asn Thr Met Cys Lys Ile Ser Gly Leu Val Gln Gly Ile Ser
469              115                 120                 125
473 Val Ala Ala Ser Val Phe Thr Leu Val Ala Ile Ala Val Asp Arg Phe
475          130                 135                 140
479 Gln Cys Val Val Tyr Pro Phe Lys Pro Lys Leu Thr Ile Lys Thr Ala
481 145                 150          ·          155                 160
485 Phe Val Ile Ile Met Ile Ile Trp Val Leu Ala Ile Thr Ile Met Ser
487              165                 170                 175
491 Pro Ser Ala Val Met Leu His Val Gln Glu Glu Lys Tyr Tyr Arg Val
493              180                 185                 190
497 Arg Leu Asn Ser Gln Asn Lys Thr Ser Pro Val Tyr Trp Cys Arg Glu
499              195                 200                 205
503 Asp Trp Pro Asn Gln Glu Met Arg Lys Ile Tyr Thr Thr Val Leu Phe
505          210                 215                 220
509 Ala Asn Ile Tyr Leu Ala Pro Leu Ser Leu Ile Val Ile Met Tyr Gly
511 225                 230                 235                 240
515 Arg Ile Gly Ile Ser Leu Phe Arg Ala Ala Val Pro His Thr Gly Arg
517              245                 250                 255
521 Lys Asn Gln Glu Gln Trp His Val Val Ser Arg Lys Lys Gln Lys Ile
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RAW SEQUENCE LISTING   DATE: 07/16/2003
PATENT APPLICATION: **US/09/866,248C**   TIME: 08:05:33

Input Set : **A:\57155A.txt**
Output Set: **N:\CRF4\07162003\I866248C.raw**

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527 Ile Lys Met Leu Leu Ile Val Ala Leu Leu Phe Ile Leu Ser Trp Leu
529           275              280              285
533 Pro Leu Trp Thr Leu Met Met Leu Ser Asp Tyr Ala Asp Leu Ser Pro
535      290              295              300
539 Asn Glu Leu Gln Ile Ile Asn Ile Tyr Ile Tyr Pro Phe Ala His Trp
541 305              310              315              320
545 Leu Ala Phe Gly Asn Ser Ser Val Asn Pro Ile Ile Tyr Gly Phe Phe
547              325              330              335
551 Asn Glu Asn Phe Arg Arg Gly Phe Gln Glu Ala Phe Gln Leu Gln Leu
553           340              345              350
557 Cys Gln Lys Arg Ala Lys Pro Met Glu Ala Tyr Ala Leu Lys Ala Lys
559           355              360              365
563 Ser His Val Leu Ile Asn Thr Ser Asn Gln Leu Val Gln Glu Ser Thr
565      370              375              380
569 Phe Gln Asn Pro His Gly Glu Thr Leu Leu Tyr Arg Lys Ser Ala Glu
571 385              390              395              400
575 Lys Pro Gln Gln Glu Leu Val Met Glu Glu Leu Lys Glu Thr Thr Asn
577           405              410              415
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583           420
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591 <211> LENGTH: 1293
593 <212> TYPE: DNA
595 <213> ORGANISM: Homo sapiens
599 <400> SEQUENCE: 7
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605 cctgtggcgg ccatgttcat tgtggcctat gcgctcatct tcctgctctg catggtgggc 180
607 aacaccctgg tctgtttcat cgtgctcaag aaccggcaca tgcatactgt caccaacatg 240
609 ttcatcctca acctggctgt cagtgacctg ctggtgggca tcttctgcat gcccaccacc 300
611 cttgtggaca acctcatcac tgggtggccc ttcgacaatg ccacatgcaa gatgagcggc 360
613 ttggtgcagg gcatgtctgt gtcggcttcc gttttcacac tggtggccat tgctgtggaa 420
615 aggttccgct gcatcgtgca ccctttccgc gagaagctga ccctgcggaa ggcgctcgtc 480
617 accatcgccg tcatctgggc cctggcgctg ctcatcatgt gtccctcggc cgtcacgctg 540
619 accgtcaccc gtgaggagca ccacttcatg gtggacgccc gcaaccgctc ctaccctctc 600
621 tactcctgct gggaggcctg gcccgagaag ggcatgcgca gggtctacac cactgtgctc 660
623 ttctcgcaca tctacctggc gccgctggcg ctcatcgtgg tcatgtacgc ccgcatcgcg 720
625 cgcaagctct gccaggcccc gggcccggcc cccggggggcg aggaggctgc ggacccgcga 780
627 gcatcgcggc gcagagcgcg cgtggtgcac atgctggtca tggtggcgct gttcttcacg 840
629 ctgtcctggc tgccgctctg ggcgctgctg ctgctcatcg actacgggca gctcagcgcg 900
631 ccgcagctgc acctggtcac cgtctacgcc ttccccttcg cgcactggct ggccttcttc 960
633 aacagcagcg ccaacccccat catctacggc tacttcaacg agaacttccg ccgcggcttc 1020
635 caggccgcct tccgcgcccg cctctgcccg cgcccgtcgg ggagccacaa ggaggcctac 1080
637 tccgagcggc ccggcggggct tctgcacagg cgggtcttcg tggtggtgcg gcccagcgac 1140
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641 ctgcggaatg ggcgggtggc tcaccacggc ttgcccaggg aagggcctgg ctgctcccac 1260
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RAW SEQUENCE LISTING ERROR SUMMARY   DATE: 07/16/2003
PATENT APPLICATION: US/09/866,248C   TIME: 08:05:34

Input Set : A:\57155A.txt
Output Set: N:\CRF4\07162003\I866248C.raw

Please Note:
Use of n and/or Xaa have been detected in the Sequence Listing. Please review the
Sequence Listing to ensure that a corresponding explanation is presented in the <220>
to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:9; N Pos. 3,9,10,12,15,21
Seq#:10; N Pos. 3,6,14,15,17,18

Invalid Line Length:
The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:1; Line(s) 1,3,5,7,9,11,13,15,17,19,21,23,25,27,29,31,33,35,37,39,41
Seq#:1; Line(s) 43,45,47,49,51,53,55,57,59,61,63,65,67,69,71,73,75,77,79,81
Seq#:1; Line(s) 83,85,87,89,91,93,95,97,99,101,103,105,107
Seq#:2; Line(s) 109,111,113,115,117,119,121,123,125,127,129,131,133,135,137
Seq#:2; Line(s) 139,141,143,145,147,149,151,153,155,157,159,161,163,165,167
Seq#:2; Line(s) 169,171,173,175,177,179,181,183,185,187,189,191,193,195,197
Seq#:2; Line(s) 199,201,203,205,207,209,211,213,215,217,219,221,223,225,227
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Seq#:2; Line(s) 259,261,263,265,267,269,271,273,275,277,279,281,283,285,287
Seq#:2; Line(s) 289
Seq#:3; Line(s) 291,293,295,297,299,301,303,305,307,309,311
Seq#:4; Line(s) 313,315,317,319,321,323,325,327,329,331,333,335,337,339,341
Seq#:4; Line(s) 343,345,347,349,351,353,355
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Seq#:5; Line(s) 387,389,391,393,395,397,399,401,403,405,407,409,411,413
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RAW SEQUENCE LISTING ERROR SUMMARY   DATE: 07/16/2003
PATENT APPLICATION: **US/09/866,248C**   TIME: 08:05:34

Input Set : **A:\57155A.txt**
Output Set: **N:\CRF4\07162003\I866248C.raw**

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Seq#:20; Line(s) 1101
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